```
   1  MGSLFQEAEP QAGTEQNKPT LASRFQQTLG DLLARLGSRG HVYVIHCLNP
  51  TPGKIPGLLD VGHVAEQLRQ AGILEIIGTR STHFPVRVSF QVFLARFHAL
 101  GSGRQKAASD QERCGAILSE VLGAESPLYH LGVTQVLLQE QGWQQLEQLW
 151  AQRRSQALLT LHRGLRACIT RQRLRLLPRM QARVRGLQAR KRYLQRRSAL
 201  GQLNTILLVA RPLLRRRQKL RCAPGPHSGE PWGKVSNMDL GRLEIPAQLA
 251  TLLERAEGHQ ALLTGSITES LPPEVPARPS LTLPPDIDQF PFSSFVSTSF
 301  QKPFLPRPGQ PLDEPLTRLD GENPQQALEI NRVMLRLLGE GSLQSWQEQT
 351  MGTFLVQQAQ RRPGLRDELF SQLVAQLWRN PDEQQNQRGW ALMVILLSSF
 401  APTPALEKPL LKFVSDQAPS GMAALCQHKL LGALEQTPLA PMASRSHPPT
 451  QLEWKAGLRR GRMALDVFTF NEESYSAEVE SWTTGEQFAG WILQSRGLEA
 501  PPRGWSVSLH SGDAWRDLPG CDFVLDLIGQ TEDLGDPAGP HNYPITPLGL
 551  AESIPPAPGV QAPSLPPGLP PGPAPILASS RPPGEASKPE NLDGFVDHLF
 601  EPALAPGFSD LEQGWALSRR MKGGGSVGPT QQGYPMVYPG MVQAPSYQPA
 651  MIPAPMPVMP AMGAVPTMPA MMVPPQPQPL VPSLDSRQLA LQQQNFINQQ
 701  AMILAQQMTT QAMSLSLEQQ NQRHQHQAQT SGATSQPPPS TTAPKAKKPP
 751  APQEKPESNL EPSGVGLRED TPEEAESKPQ RPKSFQQKRD YFQKMGQDPI
 801  RVKTVKPPAK VQIPQEEMEE TEEEEDETAE LSPPPPPPPV VKKPLKASRP
 851  KAVKEDEAEP AQEEVPTQGE DPPVHSSNSA PQHPKPSRVP PVQSSNSAPP
 901  RPQPSREIRN IIRMYQSRPG PVAVPVQPTR PIKTFQKKND PKDEALAKLG
 951  INGVHLPLST SPNQGKSSPP AVVPRPKARP RLEPSLSIQE KQGPLRDLFG
1001  PCSPNPPTAP APPPPPALPP PLSGEPKTPS VESHALTEPM EDKNISTKLL
1051  VPSGSVCFSY ANAPWKLFLR KEVFYPRENF SHPYCLSLLC QQILRDTFTE
1101  SCTRISQDER HKMKGLLGDL EVSLETLDIV EDSIKKRIVV AARDNWANYF
```

FIG. 1A.

1151 SRIFPVSGES GSDVQLLGVS HRGLRLLKVT QSPSFHLDQL KTLCSYSYAE

1201 VLTVQCRGRS TLELSLKNEQ LILHTAWARA IKAMVDLFLS ELRKDSGYVI

1251 ALRSYITDDN SLLSFHRGDL IRLLPVTALE PGWQFGSAGG RSGLFPDDVV

1301 QPAAAPDLSF SLGKRNSWQR KSKLGPAQEV RKTEEVK*

FIG. 1B.

```
   1  CGCTGGGACT GTCACCTACC AGGTGCACAA GTTCATAAAC AGAAACAGGG
  51  GCCACCTGGA CCCCGCTGTG CTGGAGATGC TCAGGCAGAG CCAGCTGCAG
 101  GTGACCTAGC CTTCCTTTCA GCTCATGGGC AGCCTGTTCC AAGAAGCAGA
 151  GCCCCAGGCT GGGACTGAGC AAAACAAACC CACATTGGCC TCTCGATTCC
 201  AGCAGACCCT GGGTGACTTG CTAGCTCGGC TAGGCAGCAG GGGCCATGTC
 251  TACGTCATCC ACTGTCTCAA TCCCACCCCT GGAAAGATCC CAGGCCTCTT
 301  GGACGTGGGG CATGTGGCAG AGCAGCTGCG TCAGGCTGGC ATCCTGGAGA
 351  TCATAGGCAC CCGGAGTACC CACTTCCCCG TGCGAGTGTC CTTCCAAGTC
 401  TTTCTGGCAA GGTTCCATGC CCTGGGGTCA GGGAGACAGA AAGCTGCCTC
 451  TGACCAGGAG AGGTGTGGTG CCATCCTCAG TGAAGTGCTG GGGGCAGAGT
 501  CACCGCTGTA TCATCTTGGA GTCACCCAGG TCCTGCTGCA GGAACAGGGC
 551  TGGCAGCAGC TAGAACAGCT GTGGGCTCAG CGGCGCTCAC AGGCCCTGCT
 601  CACTCTGCAC CGTGGCCTCC GAGCCTGTAT CACCCGGCAG CGCCTCCGTC
 651  TCCTGCCCCG GATGCAGGCT CGTGTGCGTG GGCTCCAGGC CAGGAAGCGA
 701  TATCTCCAGC GGAGGTCAGC TCTGGGACAG CTGAACACCA TTCTCCTAGT
 751  GGCCCGGCCC CTGCTCCGGA GACGACAGAA GCTACGGTGT GCCCCTGGCC
 801  CGCACAGCGG GGAGCCCTGG GGGAAAGTGT CAAATATGGA CCTGGGTCGC
 851  TTAGAGATCC CCGCCCAGCT GGCTACTCTG CTGGAGAGGG CGGAAGGCCA
 901  CCAGGCCTTG CTGACGGGGA GCATCACAGA GTCCCTGCCA CCTGAGGTCC
 951  CCGCCCGGCC CAGCCTGACT CTCCCTCCAG ACATTGACCA GTTTCCCTTC
1001  TCCAGTTTTG TATCCACCAG CTTTCAGAAG CCATTTCTGC CTCGACCAGG
1051  GCAGCCACTG GACGAGCCCC TGACGCGGTT AGATGGCGAG AACCCTCAGC
```

FIG. 2A.

```
1101  AGGCTCTGGA GATCAACAGG GTGATGCTGC GGCTCCTGGG GGAAGGATCT
1151  CTGCAGTCCT GGCAAGAGCA GACCATGGGC ACGTTCCTCG TGCAGCAGGC
1201  CCAGCGACGG CCGGGACTCC GAGATGAGCT CTTCAGCCAG CTGGTGGCCC
1251  AGCTGTGGCG CAACCCAGAT GAGCAACAGA ATCAGCGTGG CTGGGCCCTA
1301  ATGGTGATCC TGCTCAGCTC CTTTGCTCCC ACACCTGCCC TGGAGAAGCC
1351  ACTGCTCAAA TTTGTATCTG ACCAGGCTCC CAGTGGCATG GCAGCCCTGT
1401  GCCAGCACAA GCTGTTAGGT GCCCTGGAGC AGACACCGCT GGCTCCCATG
1451  GCTTCGAGGT CCCACCCACC CACACAACTT GAGTGGAAGG CTGGTTTACG
1501  TCGGGGCCGC ATGGCGCTGG ATGTGTTCAC ATTCAACGAG GAAAGCTACT
1551  CCGCGGAAGT GGAATCCTGG ACCACGGGAG AGCAGTTTGC AGGGTGGATC
1601  CTACAGAGCA GAGGCCTGGA GGCGCCCCCT CGTGGCTGGT CTGTGTCACT
1651  GCATTCTGGG GATGCTTGGC GTGACTTGCC TGGCTGTGAC TTTGTGTTGG
1701  ACCTAATAGG CCAGACTGAG GACTTGGGAG ACCCAGCTGG TCCCCACAAC
1751  TACCCCATCA CTCCTCTTGG TTTAGCTGAG AGCATCCCTC CAGCCCCTGG
1801  TGTCCAGGCT CCTTCCCTGC CCCCAGGACT CCCTCCAGGT CCAGCCCCAA
1851  TACTGGCCAG CAGCCGCCCT CCGGGCGAGG CCAGTAAGCC TGAGAACCTG
1901  GATGGTTTCG TGGACCACCT CTTTGAACCA GCGCTCGCTC CGGGTTTCAG
1951  TGATCTGGAA CAAGGCTGGG CCCTGAGCAG ACGCATGAAG GGAGGGGGCT
2001  CTGTTGGGCC CACCCAGCAG GGCTACCCCA TGGTGTACCC AGGTATGGTG
2051  CAGGCACCTA GCTACCAGCC AGCTATGATA CCCGCACCGA TGCCCGTCAT
2101  GCCAGCCATG GGCGCAGTCC CAACCATGCC AGCCATGATG GTGCCACCCC
2151  AGCCACAGCC TCTGGTGCCC AGTTTGGACT CAAGGCAGCT GGCACTACAG
2201  CAGCAAAACT TCATCAACCA GCAGGCGATG ATTCTGGCGC AGCAGATGAC
2251  CACCCAGGCC ATGAGCCTGT CCCTGGAGCA GCAGAATCAG AGACACCAGC
```

FIG. 2B.

2301 ACCAAGCTCA GACCTCTGGG GCCACCTCCC AGCCTCCACC CTCAACCACT

2351 GCTCCCAAGG CCAAGAAGCC TCCTGCCCCC CAAGAGAAGC CAGAGAGTAA

2401 CCTAGAGCCT TCGGGTGTTG GCTTGAGAGA GGACACCCCA GAGGAAGCTG

2451 AAAGCAAGCC TCAGCGCCCC AAGAGCTTCC AACAGAAACG GGACTATTTC

2501 CAGAAGATGG GGCAAGATCC GATCAGAGTG AAGACGGTGA AACCTCCAGC

2551 CAAGGTTCAG ATCCCCCAAG AGGAGATGGA GGAGACGGAG GAGGAGGAGG

2601 ATGAGACCGC CGAGTTGTCC CCTCCTCCTC CCCCTCCCCC GGTTGTGAAG

2651 AAGCCGCTGA AGGCAAGCAG GCCCAAAGCC GTAAAGGAAG ATGAGGCAGA

2701 GCCCGCCCAG GAGGAAGTAC CGACCCAGGG CGAGGATCCC CCGGTGCACA

2751 GCTCCAACTC CGCACCTCAG CACCCCAAAC CCAGCAGGGT ACCCCCAGTG

2801 CAGAGCTCCA ACTCCGCACC TCCACGCCCG CAACCCAGCA GGGAAATCCG

2851 AAACATCATC CGAATGTACC AGAGCCGTCC AGGGCCTGTG GCTGTGCCCG

2901 TACAACCCAC CAGGCCCATC AAAACTTTTC AGAAGAAAAA TGACCCTAAG

2951 GATGAGGCTT TGGCTAAGTT AGGGATAAAT GGCGTCCACT TGCCCCTATC

3001 GACATCGCCT AACCAAGGGA AGAGCTCTCC ACCGGCTGTA GTTCCTCGAC

3051 CTAAGGCTCG ACCTCGTCTT GAGCCTTCCC TATCCATCCA GGAAAAGCAG

3101 GGACCCCTTC GGGACTTGTT TGGCCCATGT AGTCCAAACC CACCTACAGC

3151 TCCAGCACCC CCGCCTCCAC CAGCACTCCC ACCGCCTCTG TCTGGGGAGC

3201 CCAAGACCCC TTCAGTGGAG TCTCATGCCT TGACAGAGCC CATGGAGGAC

3251 AAGAACATCT CCACAAAGCT CCTTGTGCCC TCTGGAAGTG TGTGCTTCTC

3301 CTATGCCAAT GCACCCTGGA AGTTGTTCTT ACGCAAGGAG GTGTTCTACC

3351 CCCGGGAGAA CTTCAGTCAT CCATACTGCC TCAGTCTCCT CTGCCAGCAG

3401 ATCCTGCGGG ACACCTTCAC AGAGTCCTGC ACCCGGATCT CACAGGATGA

FIG. 2C.

3451 GCGGCACAAA ATGAAAGGCC TTCTGGGAGA CTTGGAGGTG AGTCTGGAGA

3501 CCCTTGACAT TGTTGAAGAC AGCATCAAAA AACGCATCGT GGTCGCTGCT

3551 CGGGACAACT GGGCCAATTA CTTCTCCCGC ATCTTCCCAG TCTCGGGTGA

3601 GAGTGGCAGC GATGTACAGC TGCTGGGTGT GTCTCACCGG GGACTGCGGC

3651 TGCTGAAGGT GACCCAAAGC CCGAGCTTCC ACCTGGACCA GCTGAAGACA

3701 CTCTGTTCCT ACAGCTATGC TGAAGTCCTG ACCGTGCAGT GCAGGGGCAG

3751 ATCCACCCTG GAGCTGTCCT TGAAGAATGA GCAGCTGATA CTGCACACAG

3801 CCTGGGCGAG GGCCATCAAG GCCATGGTGG ATCTATTTCT GAGTGAACTC

3851 AGGAAGGACT CCGGCTATGT CATCGCCCTG CGCAGCTACA TCACCGATGA

3901 CAATAGCCTC CTCAGTTTCC ACCGTGGGGA CCTCATTAGG TTACTGCCAG

3951 TGACCGCTCT GGAACCAGGC TGGCAGTTCG GTTCTGCCGG GGGCCGCTCC

4001 GGACTCTTTC CCGATGACGT GGTGCAGCCA GCTGCTGCCC CCGACCTCTC

4051 CTTTTCCCTG GGAAAGAGAA ACAGCTGGCA ACGCAAGAGT AAGCTGGGGC

4101 CAGCTCAGGA GGTGAGGAAG ACAGAAGAGG TGAAGTGATA CAGGCCTAAC

4151 TTGGAGACTG AGAAGGAAAG AGCAGGGTTG CTTCGGGTGT TGTCCACTTC

4201 CTGTCCTGGT GGCCAGGGCT CAATGTGTTC CTGTCCTTTA CCATCTCCTG

4251 ACTTTTTGCC ATTTGTGAGA CTGTAAGTCA CACCCTCTAA CTCTGGTACT

4301 TAGTTCAGTG TCTCCATAGA GGATGCTTAA TAAATAACCT TGGTTTTCCT

4351 GGTTTCTGGT GTCACTCCTC TTGGGTCTAA TGGGTATGGG GACCAGGGCC

4401 TGAGAGTGAG TATTGGGCCT CTGGGCTAGA TGGTGGGTAC TGGGGTGGTA

4451 CCAAATTTCC TGTGCTCCCA GCGCCCCACC CATCCCAGGA AACAAGAACC

4501 CAGTGAAGAC TCGGAGGCCA CCTCCTTTAC AACCTACAGC TCTTTGTCTG

4551 CCGACCCCCA CAACTACACC ATGCAGGAAT TTGCCCTGCG CTATTTCCGG

4601 AAGCCTCATA CCTGGCTGAC CCAGATGAGT AGAGACACCA AAGAGAAAGC

FIG. 2D.

```
4651  TGCCATCAAC CTGATCCAGT ACACTAAGGA CCCCATCCAG GAATCCCTTA
4701  CCAGCTTCTG CAATGGGGAC ACAAACAGTA AAGCTGTGGC TGGCTTCAAG
4751  GCTCTGATGC AGTTTATGGG GGACCAGCCT AAGCCCCGGG GCAAGGACGA
4801  GCTGAGTCTG CTCTATGAGC TGCTGAAGCT GTGCCAAGAT GACCTTAGGG
4851  ACGAGATGTA CTGCCAGGTC ATCAAGCAAG TCACAGGACA CCCCCAGCCA
4901  AAGCACTGTG CTCTGGGCTG GAGCGTCCTC AGCCTCTTCA CAGGCTTCTT
4951  TGCACCATCG ACCACGCTGA TGCCCTATGT GACCAAGTTC CTGCAGGATT
5001  CCAGCCCCAG TGAAGAGTTG GCCAGGAGGA GCCAGGAGAA CCTCCAGCGC
5051  ACAGTTAAAT ATGGGGGACG CCAGCAGCTG CCGTTACCTG GTGAAATGAA
5101  TGCTTTTCTG AAAGGGCAAG CAGTTCGTTT GCTTCTAATT CACCTGCCTG
5151  GGGGTGTGGA CTACAGGACG AATTCACAGA CATTCACAGT GGCAGGGGAA
5201  GTGCTAGAGG AGCTGTGTGG ACAGATGGGC ATCACAGACT TGGAAGAAGT
5251  GCAGGAATTT GCCCTCTTTC TCATCAAAGG AGAAGGTGAG CTGGTTCGGC
5301  CGCTGTCACC CCATGAGTAC ATCAACAATG TGGTGACGGA CCAGGACATG
5351  AGCCTTCACA GCCGACGGCT TGGTTGGGAG ACTCCACTGC ATTTTGATCA
5401  CTCCACCTAC ACGGAAACCC ACTATGGCCA GGTGCTTCGG GACTACCTGC
5451  AAGGGAAGCT GATAGTCAGC ACCCAGGCAG AGGCTCTACT TGCCCAGCTT
5501  GCTGCCTTCC AACACTTCGA CAAAACCGGA ACTTCTAGTC CTCCATCAGA
5551  GCAAGAGCTG CTGTCTTATA TTCCCAAGCC ACTGCAATGG CAGGTGAACA
5601  CAGCCAACAT AAAGAGCTTG GTGACCCAGG AGCTGAGGCA GATGCAAGGG
5651  TACAGCAAGC AGAGAGCACA GATTGGCTTT ATAGAGAGCA CAGCGCAGCT
5701  GCCCCTCTTT GGCTACACTG TGTACGTAGT GCTGAGAGTG AGTAAGCTGG
5751  CCCTCCCTGG ACCAGGCCTC CTGGGGCTGA ACCGTCAGCA CCTGGTCCTC
```

FIG. 2E.

5801 ATGGACCCCA GCTCTCAGGA ACTCTGCTGC TCTGTCATGC TAAAAGACCT

5851 GAAGCAGTTC CACCTGCTGA GCCCACTGCA GGAGGACGGG CCCCCTGGCC

5901 TAGAACTCAA CTATGGCTCT GTTGACAACC CCCAGACCAT CTGGTTGGAG

5951 TTGCCACAGG CCCAGGAGCT GCAGCACACC ATCATCTTCC TGCTGGGCAG

6001 CATGTCCACT CAGTGGCCAG GTCTCCTCTG AGGAGTGGAG ATAAGGCAGC

6051 GGTCTCTCAC TGGGCAGTCT GCCTTAGTCC TGCTCTGAAT CCGCTGCACA

6101 ACCCCCCACC CCACGTGGAG GCCAAAAGGC AAAGTTGTGT CACCTGGGAG

6151 AATAGGCAGA CACATCCCCT CTGGGGTGGA CTGCAACAGG AGTTGGGGCA

6201 TTTGCTGGCT AGCCCCAGGG AAAATGCCCA CCCAGCTCGA AAGCGGCACA

6251 AGTAAAACAC CCAAGGAAAA AAAAAAAAAA AAAAAAAAAA AAA

FIG. 2F.

```
   1  CGCTGGGACT GTCACCTACC AGGTGCACAA GTTCATAAAC AGAAACAGGG
  51  GCCACCTGGA CCCCGCTGTG CTGGAGATGC TCAGGCAGAG CCAGCTGCAG
 101  GTGACCTAGC CTTCCTTTCA GCTCATGGGC AGCCTGTTCC AAGAAGCAGA
 151  GCCCCAGGCT GGGACTGAGC AAAACAAACC CACATTGGCC TCTCGATTCC
 201  AGCAGACCCT GGGTGACTTG CTAGCTCGGC TAGGCAGCAG GGGCCATGTC
 251  TACGTCATCC ACTGTCTCAA TCCCACCCCT GGAAAGATCC CAGGCCTCTT
 301  GGACGTGGGG CATGTGGCAG AGCAGCTGCG TCAGGCTGGC ATCCTGGAGA
 351  TCATAGGCAC CCGGAGTACC CACTTCCCCG TGCGAGTGTC CTTCCAAGTC
 401  TTTCTGGCAA GGTTCCATGC CCTGGGGTCA GGGAGACAGA AAGCTGCCTC
 451  TGACCAGGAG AGGTGTGGTG CCATCCTCAG TGAAGTGCTG GGGGCAGAGT
 501  CACCGCTGTA TCATCTTGGA GTCACCCAGG TCCTGCTGCA GGAACAGGGC
 551  TGGCAGCAGC TAGAACAGCT GTGGGCTCAG CGGCGCTCAC AGGCCCTGCT
 601  CACTCTGCAC CGTGGCCTCC GAGCCTGTAT CACCCGGCAG CGCCTCCGTC
 651  TCCTGCCCCG GATGCAGGCT CGTGTGCGTG GGCTCCAGGC CAGGAAGCGA
 701  TATCTCCAGC GGAGGTCAGC TCTGGGACAG CTGAACACCA TTCTCCTAGT
 751  GGCCCGGCCC CTGCTCCGGA GACGACAGAA GCTACGGTGT GCCCCTGGCC
 801  CGCACAGCGG GGAGCCCTGG GGGAAAGTGT CAAATATGGA CCTGGGTCGC
 851  TTAGAGATCC CCGCCCAGCT GGCTACTCTG CTGGAGAGGG CGGAAGGCCA
 901  CCAGGCCTTG CTGACGGGGA GCATCACAGA GTCCCTGCCA CCTGAGGTCC
 951  CCGCCCGGCC CAGCCTGACT CTCCCTCCAG ACATTGACCA GTTTCCCTTC
1001  TCCAGTTTTG TATCCACCAG CTTTCAGAAG CCATTTCTGC CTCGACCAGG
1051  GCAGCCACTG GACGAGCCCC TGACGCGGTT AGATGGCGAG AACCCTCAGC
```

FIG. 3A.

1101 AGGCTCTGGA GATCAACAGG GTGATGCTGC GGCTCCTGGG GGAAGGATCT

1151 CTGCAGTCCT GGCAAGAGCA GACCATGGGC ACGTTCCTCG TGCAGCAGGC

1201 CCAGCGACGG CCGGGACTCC GAGATGAGCT CTTCAGCCAG CTGGTGGCCC

1251 AGCTGTGGCG CAACCCAGAT GAGCAACAGA ATCAGCGTGG CTGGGCCCTA

1301 ATGGTGATCC TGCTCAGCTC CTTTGCTCCC ACACCTGCCC TGGAGAAGCC

1351 ACTGCTCAAA TTTGTATCTG ACCAGGCTCC CAGTGGCATG GCAGCCCTGT

1401 GCCAGCACAA GCTGTTAGGT GCCCTGGAGC AGACACCGCT GGCTCCCATG

1451 GCTTCGAGGT CCCACCCACC CACACAACTT GAGTGGAAGG CTGGTTTACG

1501 TCGGGGCCGC ATGGCGCTGG ATGTGTTCAC ATTCAACGAG GAAAGCTACT

1551 CCGCGGAAGT GGAATCCTGG ACCACGGGAG AGCAGTTTGC AGGGTGGATC

1601 CTACAGAGCA GAGGCCTGGA GGCGCCCCCT CGTGGCTGGT CTGTGTCACT

1651 GCATTCTGGG GATGCTTGGC GTGACTTGCC TGGCTGTGAC TTTGTGTTGG

1701 ACCTAATAGG CCAGACTGAG GACTTGGGAG ACCCAGCTGG TCCCCACAAC

1751 TACCCCATCA CTCCTCTTGG TTTAGCTGAG AGCATCCCTC CAGCCCCTGG

1801 TGTCCAGGCT CCTTCCCTGC CCCCAGGACT CCCTCCAGGT CCAGCCCCAA

1851 TACTGGCCAG CAGCCGCCCT CCGGGCGAGG CCAGTAAGCC TGAGAACCTG

1901 GATGGTTTCG TGGACCACCT CTTTGAACCA GCGCTCGCTC CGGGTTTCAG

1951 TGATCTGGAA CAAGGCTGGG CCCTGAGCAG ACGCATGAAG GGAGGGGGCT

2001 CTGTTGGGCC CACCCAGCAG GGCTACCCCA TGGTGTACCC AGGTATGGTG

2051 CAGGCACCTA GCTACCAGCC AGCTATGATA CCCGCACCGA TGCCCGTCAT

2101 GCCAGCCATG GGCGCAGTCC CAACCATGCC AGCCATGATG GTGCCACCCC

2151 AGCCACAGCC TCTGGTGCCC AGTTTGGACT CAAGGCAGCT GGCACTACAG

2201 CAGCAAAACT TCATCAACCA GCAGGCGATG ATTCTGGCGC AGCAGATGAC

2251 CACCCAGGCC ATGAGCCTGT CCCTGGAGCA GCAGAATCAG AGACACCAGC

FIG. 3B.

2301 ACCAAGCTCA GACCTCTGGG GCCACCTCCC AGCCTCCACC CTCAACCACT

2351 GCTCCCAAGG CCAAGAAGCC TCCTGCCCCC CAAGAGAAGC CAGAGAGTAA

2401 CCTAGAGCCT TCGGGTGTTG GCTTGAGAGA GGACACCCCA GAGGAAGCTG

2451 AAAGCAAGCC TCAGCGCCCC AAGAGCTTCC AACAGAAACG GGACTATTTC

2501 CAGAAGATGG GGCAAGATCC GATCAGAGTG AAGACGGTGA AACCTCCAGC

2551 CAAGGTTCAG ATCCCCCAAG AGGAGATGGA GGAGACGGAG GAGGAGGAGG

2601 ATGAGACCGC CGAGTTGTCC CCTCCTCCTC CCCCTCCCCC GGTTGTGAAG

2651 AAGCCGCTGA AGGCAAGCAG GCCCAAAGCC GTAAAGGAAG ATGAGGCAGA

2701 GCCCGCCCAG GAGGAAGTAC CGACCCAGGG CGAGGATCCC CCGGTGCACA

2751 GCTCCAACTC CGCACCTCAG CACCCCAAAC CCAGCAGGGT ACCCCCAGTG

2801 CAGAGCTCCA ACTCCGCACC TCCACGCCCG CAACCCAGCA GGGAAATCCG

2851 AAACATCATC CGAATGTACC AGAGCCGTCC AGGGCCTGTG GCTGTGCCCG

2901 TACAACCCAC CAGGCCCATC AAAACTTTTC AGAAGAAAAA TGACCCTAAG

2951 GATGAGGCTT TGGCTAAGTT AGGGATAAAT GGCGTCCACT TGCCCCTATC

3001 GACATCGCCT AACCAAGGGA AGAGCTCTCC ACCGGCTGTA GTTCCTCGAC

3051 CTAAGGCTCG ACCTCGTCTT GAGCCTTCCC TATCCATCCA GGAAAAGCAG

3101 GGACCCCTTC GGGACTTGTT TGGCCCATGT AGTCCAAACC CACCTACAGC

3151 TCCAGCACCC CCGCCTCCAC CAGCACTCCC ACCGCCTCTG TCTGGGGAGC

3201 CCAAGACCCC TTCAGTGGAG TCTCATGCCT TGACAGAGCC CATGGAGGAC

3251 AAGAACATCT CCACAAAGCT CCTTGTGCCC TCTGGAAGTG TGTGCTTCTC

3301 CTATGCCAAT GCACCCTGGA AGTTGTTCTT ACGCAAGGAG GTGTTCTACC

3351 CCCGGGAGAA CTTCAGTCAT CCATACTGCC TCAGTCTCCT CTGCCAGCAG

3401 ATCCTGCGGG ACACCTTCAC AGAGTCCTGC ACCCGGATCT CACAGGATGA

FIG. 3C.

3451 GCGGCACAAA ATGAAAGGCC TTCTGGGAGA CTTGGAGGTG AGTCTGGAGA

3501 CCCTTGACAT TGTTGAAGAC AGCATCAAAA AACGCATCGT GGTCGCTGCT

3551 CGGGACAACT GGGCCAATTA CTTCTCCCGC ATCTTCCCAG TCTCGGGTGA

3601 GAGTGGCAGC GATGTACAGC TGCTGGGTGT GTCTCACCGG GGACTGCGGC

3651 TGCTGAAGGT GACCCAAAGC CCGAGCTTCC ACCTGGACCA GCTGAAGACA

3701 CTCTGTTCCT ACAGCTATGC TGAAGTCCTG ACCGTGCAGT GCAGGGGCAG

3751 ATCCACCCTG GAGCTGTCCT TGAAGAATGA GCAGCTGATA CTGCACACAG

3801 CCTGGGCGAG GGCCATCAAG GCCATGGTGG ATCTATTTCT GAGTGAACTC

3851 AGGAAGGACT CCGGCTATGT CATCGCCCTG CGCAGCTACA TCACCGATGA

3901 CAATAGCCTC CTCAGTTTCC ACCGTGGGGA CCTCATTAGG TTACTGCCAG

3951 TGACCGCTCT GGAACCAGGC TGGCAGTTCG GTTCTGCCGG GGGCCGCTCC

4001 GGACTCTTTC CCGATGACGT GGTGCAGCCA GCTGCTGCCC CCGACCTCTC

4051 CTTTTCCCTG GGAAAGAGAA ACAGCTGGCA ACGCAAGAGT AAGCTGGGGC

4101 CAGCTCAGGA GGTGAGGAAG ACAGAAGAGG TGAAGTGATA CAGGCCTAAC

4151 TTGGAGACTG AGAAGGAAAG AGCAGGGTTG CTTCGGGTGT TGTCCACTTC

4201 CTGTCCTGGT GGCCAGGGCT CAATGTGTTC CTGTCCTTTA CCATCTCCTG

4251 ACTTTTTGCC ATTTGTGAGA CTGTAAGTCA CACCCTCTAA CTCTGGTACT

4301 TAGTTCAGTG TCTCCATAGA GGATGCTTAA TAAATAACCT TGGTTTTCCT

4351 GGAAAAAAAA AAAAAAAAAA AAAAA

FIG. 3D.

MYQSRPGPVPVPVQPSRPPKAFLRKIDPKDEALAKLGINGAHSSPPMLSPSPGKGPPPAVAPRPKA
PLQLGPSSSIKEKQGPLLDLFGQKLPIAHTPPPPPAPPLPLPEDPGTLSAERRCLTQPVEDQGVST
QLLAPSGSVCFSYTGTPWKLFLRKEVFYPRENFSHPYYLRLLCEQILRDTFSESCIRISQNERRKM
KDLLGGLEVDLDSLTTTEDSVKKRIVVAARDNWANYFSRFFPVSGESGSDVQLLAVSHRGLRLLKV
TQGPGLRPDQLKILCSYSFAEVLGVECRGGSTLELSLKSEQLVLHTARARAIEALVELFLNELKKD
SGYVIALRSYITDNCSLLSFHRGDLIKLLPVATLEPGWQFGSAGGRSGLFPADIVQPAAAPDFSFS
KEQRSGWHKGQLSNGEPGLARWDRASEVRKMGEGQAEARPA

FIG. 4.

CGGCAGCAGCAGGCTCGGGCCTCCGAGGCTGCGTCCCAGGCCTCACCCTCAGCCGTCACCTCCAAG
CCCAGGAAGCCCCCCACACCCCCGGAGAAGCCACAGCGTGACCTGGGATCAGAGGGTGGCTGCCTG
AGGGAGACCTCCGAGGAGGCTGAAGACAGGCCCTATCAGCCCAAGAGCTTCCAGCAGAAACGGAAC
TATTTCCAGAGGATGGGGCAGCCACAGATCACAGTGAGGACGATGAAGCCCCCGGCCAAGGTCCAC
ATCCCCCAGGGGGAAGCGCAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGCAGGAGGAGCAA
GAAGTGGAAACAAGAGCAGCGCCGTCCCCTCCTCCTCCCCCCATCGTGAAGAAGCCATTGAAGCAA
GGTGGGGCCAAAGCTCCAAAAGAGGCTGAGGCTGAGCCAGCCAAGGAGACAGCGGCCAAGGGCCAT
GGCCAAGGGCCAGCCCAAGGCAGGGGGACTGTGGTGCGCAGTCAGACTCCAAGCCCAAGCGGCCAC
AACCCAGCAGGGAAATTGGCAACATCATCCGCATGTACCAGAGCCGCCCGGGCCCCGTGCCTGTGC
CCGTGCAGCCATCCAGGCCTCCCAAAGCTTTCCTGAGGAAAATCGACCCCAAGGACGAGGCTCTGG
CCAAGCTGGGTATCAACGGTGCCCACTCGTCCCCGCCGATGCTGTCCCCCAGCCCAGGAAAGGGCC
CCCCGCCAGCTGTGGCTCCTCGACCCAAGGCCCCGCTACAGCTTGGGCCCTCTAGCTCCATCAAGG
AAAAGCAGGGGCCCCTTCTGGACCTGTTTGGCCAGAAGCTGCCTATTGCCCACACACCCCCACCTC
CACCAGCGCCACCACTGCCTCTGCCCGAGGACCCAGGGACCCTTTCAGCAGAGCGTCGTTGCTTGA
CACAGCCCGTGGAGGACCAGGGGGTCTCCACCCAGCTACTCGCGCCCTCTGGCAGCGTGTGCTTCT
CCTACACCGGCACGCCCTGGAAGTTGTTCCTACGCAAGGAGGTGTTCTACCCACGGGAGAACTTCA
GCCATCCCTACTACCTGAGGCTCCTCTGTGAGCAGATCCTACGGGACACCTTCTCCGAGTCCTGTA
TCCGGATTTCCCAGAATGAGCGGCGGAAAATGAAAGACCTGCTGGGAGGCTTGGAGGTGGACCTGG
ATTCTCTCACCACCACCGAAGACAGCGTCAAGAAGCGCATCGTGGTGGCCGCTCGGGACAACTGGG
CCAATTACTTCTCCCGCTTCTTTCCTGTCTCGGGCGAGAGTGGCAGCGACGTGCAGCTGTTAGCCG
TGTCCCACCGTGGGCTGCGACTGCTCAAGGTGACCCAAGGCCCCGGCCTCCGCCCCGACCAGCTGA
AGATTCTCTGCTCATACAGCTTTGCGGAGGTGCTGGGTGTGGAGTGCCGGGGCGGCTCCACCCTGG
AGCTGTCACTGAAGAGCGAGCAGCTGGTGCTGCACACAGCCCGGGCAAGGGCCATCGAGGCGCTGG
TTGAGCTATTCCTGAATGAGCTTAAGAAGGACTCCGGCTATGTCATCGCCCTGCGCAGCTACATCA
CTGACAACTGCAGCCTCCTCAGCTTCCACCGTGGGGACCTCATCAAGCTGCTGCCGGTGGCCACCC
TGGAGCCAGGCTGGCAGTTTGGCTCTGCCGGGGGCCGTTCCGGACTCTTTCCTGCCGACATAGTGC
AGCCGGCTGCCGCTCCCGACTTTTCCTTCTCCAAGGAGCAGAGGAGTGGCTGGCACAAGGGTCAGC
TGTCCAACGGGGAACCAGGGCTGGCTCGGTGGGACAGGGCCTCAGAGGTGAGGAAGATGGGAGAGG
GACAAGCAGAGGCAAGGCCTGCCTGAGACTGAGGAAGGAAAGGGGTTTGACCACTCCCGAGGCTGC
CATGCGGTGGGACCACCCTGCTGTCCGTCTCCTGTGGCTGCCCCTCTGCCCGCTCCTGATGGCTCG
CCTTGTCTCTCCAGCAAGACTGTGCACTCCTTGCAGGCAGGGGCTGGGCTGGATGCTGCTCTTGTG

FIG. 5A.

TCCCACGTGGTACTTAGTTCAAGGCTGCCCCAGCAGATGCTTAATAAACAGCTCTTCACTTTCCTG
GCTTCTGGTCTTGCTCCTTTGGTGTCTGGCTGGGGAGGGATGGGGCTGGGGCAGGACCCCTGGGAC
AGGGCACTGGACACTCAGGTGGCACCAGGTTTCTTGTGATCCCAGCGCCCTGCCCACCCTTGGAGC
CAGGCACACAGTGACGACTCGGAGGCCACCAGCCTGTCCTCTGTGGCCTATGCCTTTCTGCCCGAC
TCCCACAGCTACACCATGCAGGAATTCGCCCGGCGTTACTTCCGGAGGTCCCAGGCCTTGCTGGGC
CAGACTGATGGAGGTGCCGCAGGAAAGGACACGGACAGCCTGGTGCAGTACACCAAGGCTCCCATC
CAGGAGTCGCTCCTCAGCCTCAGTGATGATGTGAGCAAGCTGGCTGTAGCCAGCTTCCTGGCCCCT
GATGCGGTTTATGGGTGACCAGTCCAAGCCCCGGGGCAAGGATGAGATGGATCTGCTCTATGAACT
GCTGAAGCTGTGCCAGCAGGAGAAGCTGAGGGATGAGATTTACTGCCAGGTTATCAAGCAGGTCAC
AGGACACCCCCGGCCGGAACACTGCACTCGAGGCTGGAGCTTCCTCAGCCTTCTCACAGGCTTCTT
CCCCCCGTCGACCAGGCTGATGCCCTACCTGACCAAGTTTCTGCAGGATTCAGGCCCCAGCCAAGA
GCTGGCCCGGAGCAGCCAGGAGCACCTCCAGCGCACAGTCAAATATGGGGGGCGCCGGCGGATGCC
CCCACCGGGTGAAATGAAGGCTTTCCTGAAAGGACAAGCGATTCGCCTGCTTCTTATTCACCTGCC
GGGGGGTGTGGATTATAGGACGAATATCCAGACTTTCACAGTAGCAGCAGAAGTGCAGGAGGAGCT
GTGCCGGCAAATGGGTATCACGGAGCCTCAGGAAGTGCAGGAATTCGCCCTCTTCCTCATCAAAGA
GAAGAGCCAGCTGGTGCGGCCCCTGCAGCCCGCCGAATACCTCAACAGCGTGGTAGTGGACCAGGA
CGTGAGCCTGCACAGCCGGCGGCTCCACTGGGAGACCCCACTGCACTTCGATAACTCCACCTACAT
CAGCACCCACTACAGCCAGGTGCTGTGGGACTACCTTCAGGGGAAGCTGCCAGTCAGCGCCAAGGC
AGACGCGCAGCTCGCCAGGCTGGCCGCCCTGCAGCACCTCAGCAAGGCCAACACGGAATACCCCCTC
AGGGCAGGACCTGCTAGCTTACGTGCCAAAGCAGCTGCAACGGCAGGTGAACACGGCCTCCATCAA
GAACCTGATGGGTCAGGAGCTGAGACGGCTGGAAGGACACAGCCCCCAGGAAGCACAGATCAGCTT
CATTGAGGCCATGAGCCAGCTGCCCCTCTTCGGCTACACCGTCTATGGGGTGCTGCGAGTGAGCAT
GCAGGCCCTGTCCGGACCCACTCTCCTGGGGCTCAACCGCCAGCATCTCATCCTCATGGACCCCAG
CTCCCAGAGCCTGTACTGCCGCATTGCCCTGAAGAGCCTGCAGCGGCTCCACCTGCTAAGCCCTCT
GGAGGAGAAGGGGCCCCCTGGCCTGGAAGTCAACTATGGCTCAGCTGACAACCCCCAGACCATCTG
GTTTGAGCTGCCACAGGCCCAGGAGCTGCTATACACCACTGTCTTCCTGATAGACAGCAGTGCCTC
TTGCACTGAGTGGCCCAGCATCAACTGAGAGGAGTGCAGGCCGGGGAGAGAAGAGGATGAGGCCTC
CCCCGGCCCAAGTCTCACCCACATGGTCTGCCTTGGATGCTATCAGATCACTGTTCTAGAACCTGC
CTCAGCACAGCCCAGCCGGCCCACATGCAGGCCATGAGGCAGGGGCTGCTATCACGTCACCAGCAG
GCAAAGAAAACAGCCAGACCCTCTCCAGGACGGCCTGGGGCCAAAGCGGGCTGCAGGAACTCGGCT
GGGGCACCTGAGGTTGCCCAGTCTGAGGGAGATGCCCACCCGACCCCAGGCTCCGCCCAGGCCCCA

FIG. 5B.

CATTAGCACAAGCCCAGGCATGGGAGAAACAGCTGCTGAGGAAATAAAACTCCCTAAAAAAAAAAA
AAAAAAAAAAAAAAAA

FIG. 5C.

MYQSRPGPVPVPVQPSRPPKAFLRKIDPKDEALAKLGINGAHSSPPMLSPSPGKGPPPAVAPRPKA
PLQLGPSSSIKEKQGPLLDLFGQKLPIAHTPPPPPAPPLPLPEDPGTLSAERRCLTQPVEDQGVST
QLLAPSGSVCFSYTGTPWKLFLRKEVFYPRENFSHPYYLRLLCEQILRDTFSESCIRISQNERRKM
KDLLGGLEVDLDSLTTTEDSVKKRIVVAARDNWANYFSRFFPVSGESGSDVQLLAVSHRGLRLLKV
TQGPGLRPDQLKILCSYSFAEVLGVECRGGSTLELSLKSEQLVLHTARARAIEALVELFLNELKKD
SGYVIALRSYITDNCSLLSFHRGDLIKLLPVATLEPGWQFGSAGGRSGLFPADIVQPAAAPDFSFS
KEQRSGWHKGQLSNGEPGLARWDRASERPAHPWSQAHSDDSEATSLSSVAYAFLPDSHSYTMQEFA
RRYFRRSQALLGQTDGGAAGKDTDSLVQYTKAPIQESLLSLSDDVSKLAVASFLALMRFMGDQSKP
RGKDEMDLLYELLKLCQQEKLRDEIYCQVIKQVTGHPRPEHCTRGWSFLSLLTGFFPPSTRLMPYL
TKFLQDSGPSQELARSSQEHLQRTVKYGGRRRMPPPGEMKAFLKGQAIRLLLIHLPGGVDYRTNIQ
TFTVAAEVQEELCRQMGITEPQEVQEFALFLIKEKSQLVRPLQPAEYLNSVVVDQDVSLHSGGSTG
RPHCTSITPPTSAPTTARCCGTTFRGSCQSAPRQTRSSPGWPPCSTSARPTGIPPQGRTC

FIG. 6.

CGGCAGCAGCAGGCTCGGGCCTCCGAGGCTGCGTCCCAGGCCTCACCCTCAGCCGTCACCTCCAAG
CCCAGGAAGCCCCCCACACCCCCGGAGAAGCCACAGCGTGACCTGGGATCAGAGGGTGGCTGCCTG
AGGGAGACCTCCGAGGAGGCTGAAGACAGGCCCTATCAGCCCAAGAGCTTCCAGCAGAAACGGAAC
TATTTCCAGAGGATGGGGCAGCCACAGATCACAGTGAGGACGATGAAGCCCCCGGCCAAGGTCCAC
ATCCCCCAGGGGGAAGCGCAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGCAGGAGGAGCAA
GAAGTGGAAACAAGAGCAGCGCCGTCCCCTCCTCCTCCCCCCATCGTGAAGAAGCCATTGAAGCAA
GGTGGGGCCAAAGCTCCAAAAGAGGCTGAGGCTGAGCCAGCCAAGGAGACAGCGGCCAAGGGCCAT
GGCCAAGGGCCAGCCCAAGGCAGGGGGACTGTGGTGCGCAGTCAGACTCCAAGCCCAAGCGGCCAC
AACCCAGCAGGGAAATTGGCAACATCATCCGCATGTACCAGAGCCGCCCGGGCCCCGTGCCTGTGC
CCGTGCAGCCATCCAGGCCTCCCAAAGCTTTCCTGAGGAAAATCGACCCCAAGGACGAGGCTCTGG
CCAAGCTGGGTATCAACGGTGCCCACTCGTCCCCGCCGATGCTGTCCCCCAGCCCAGGAAAGGGCC
CCCCGCCAGCTGTGGCTCCTCGACCCAAGGCCCCGCTACAGCTTGGGCCCTCTAGCTCCATCAAGG
AAAAGCAGGGGCCCCTTCTGGACCTGTTTGGCCAGAAGCTGCCTATTGCCCACACACCCCCACCTC
CACCAGCGCCACCACTGCCTCTGCCCGAGGACCCAGGGACCCTTTCAGCAGAGCGTCGTTGCTTGA
CACAGCCCGTGGAGGACCAGGGGGTCTCCACCCAGCTACTCGCGCCCTCTGGCAGCGTGTGCTTCT
CCTACACCGGCACGCCCTGGAAGTTGTTCCTACGCAAGGAGGTGTTCTACCCACGGGAGAACTTCA
GCCATCCCTACTACCTGAGGCTCCTCTGTGAGCAGATCCTACGGGACACCTTCTCCGAGTCCTGTA
TCCGGATTTCCCAGAATGAGCGGCGGAAAATGAAAGACCTGCTGGGAGGCTTGGAGGTGGACCTGG
ATTCTCTCACCACCACCGAAGACAGCGTCAAGAAGCGCATCGTGGTGGCCGCTCGGGACAACTGGG
CCAATTACTTCTCCCGCTTCTTTCCTGTCTCGGGCGAGAGTGGCAGCGACGTGCAGCTGTTAGCCG
TGTCCCACCGTGGGCTGCGACTGCTCAAGGTGACCCAAGGCCCCGGCCTCCGCCCCGACCAGCTGA
AGATTCTCTGCTCATACAGCTTTGCGGAGGTGCTGGGTGTGGAGTGCCGGGGCGGCTCCACCCTGG
AGCTGTCACTGAAGAGCGAGCAGCTGGTGCTGCACACAGCCCGGGCAAGGGCCATCGAGGCGCTGG
TTGAGCTATTCCTGAATGAGCTTAAGAAGGACTCCGGCTATGTCATCGCCCTGCGCAGCTACATCA
CTGACAACTGCAGCCTCCTCAGCTTCCACCGTGGGGACCTCATCAAGCTGCTGCCGGTGGCCACCC
TGGAGCCAGGCTGGCAGTTTGGCTCTGCCGGGGGCCGTTCCGGACTCTTTCCTGCCGACATAGTGC
AGCCGGCTGCCGCTCCCGACTTTTCCTTCTCCAAGGAGCAGAGGAGTGGCTGGCACAAGGGTCAGC
TGTCCAACGGGGAACCAGGGCTGGCTCGGTGGGACAGGGCCTCAGAGCGCCCTGCCCACCCTTGGA
GCCAGGCACACAGTGACGACTCGGAGGCCACCAGCCTGTCCTCTGTGGCCTATGCCTTTCTGCCCG
ACTCCCACAGCTACACCATGCAGGAATTCGCCCGGCGTTACTTCCGGAGGTCCCAGGCCTTGCTGG
GCCAGACTGATGGAGGTGCCGCAGGAAAGGACACGGACAGCCTGGTGCAGTACACCAAGGCTCCCA

FIG. 7A.

TCCAGGAGTCGCTCCTCAGCCTCAGTGATGATGTGAGCAAGCTGGCTGTAGCCAGCTTCCTGGCCC
TGATGCGGTTTATGGGTGACCAGTCCAAGCCCCGGGGCAAGGATGAGATGGATCTGCTCTATGAAC
TGCTGAAGCTGTGCCAGCAGGAGAAGCTGAGGGATGAGATTTACTGCCAGGTTATCAAGCAGGTCA
CAGGACACCCCCGGCCGGAACACTGCACTCGAGGCTGGAGCTTCCTCAGCCTTCTCACAGGCTTCT
TCCCCCCGTCGACCAGGCTGATGCCCTACCTGACCAAGTTTCTGCAGGATTCAGGCCCCAGCCAAG
AGCTGGCCCGGAGCAGCCAGGAGCACCTCCAGCGCACAGTCAAATATGGGGGGCGCCGGCGGATGC
CCCCACCGGGGTGAAATGAAGGCTTTCCTGAAAGGACAAGCGATTCGCCTGCTTCTTATTCACCTGC
CGGGGGGTGTGGATTATAGGACGAATATCCAGACTTTCACAGTAGCAGCAGAAGTGCAGGAGGAGC
TGTGCCGGCAAATGGGTATCACGGAGCCTCAGGAAGTGCAGGAATTCGCCCTCTTCCTCATCAAAG
AGAAGAGCCAGCTGGTGCGGCCCCTGCAGCCCGCCGAATACCTCAACAGCGTGGTAGTGGACCAGG
ACGTGAGCCTGCACAGCGGCGGCTCCACTGGGAGACCCCACTGCACTTCGATAACTCCACCTACAT
CAGCACCCACTACAGCCAGGTGCTGTGGGACTACCTTCAGGGGAAGCTGCCAGTCAGCGCCAAGGC
AGACGCGCAGCTCGCCAGGCTGGCCGCCCTGCAGCACCTCAGCAAGGCCAACAGGAATACCCCCTC
AGGGCAGGACCTGCTAGCTTACGTGCCAAAGCAGCTGCAACGGCAGGTGAACACGGCCTCCATCAA
GAACCTGATGGGTCAGGAGCTGAGACGGCTGGAAGGACACAGCCCCCAGGAAGCACAGATCAGCTT
CATTGAGGCCATGAGCCAGCTGCCCCTCTTCGGCTACACCGTCTATGGGGTGCTGCGAGTGAGCAT
GCAGGCCCTGTCCGGACCCACTCTCCTGGGGCTCAACCGCCAGCATCTCATCCTCATGGACCCCAG
CTCCCAGAGCCTGTACTGCCGCATTGCCCTGAAGAGCCTGCAGCGGCTCCACCTGCTAAGCCCTCT
GGAGGAGAAGGGGCCCCCTGGCCTGGAAGTCAACTATGGCTCAGCTGACAACCCCCAGACCATCTG
GTTTGAGCTGCCACAGGCCCAGGAGCTGCTATACACCACTGTCTTCCTGATAGACAGCAGTGCCTC
TTGCACTGAGTGGCCCAGCATCAACTGAGAGGAGTGCAGGCCGGGGAGAGAAGAGGATGAGGCCTC
CCCCGGCCCAAGTCTCACCCACATGGTCTGCCTTGGATGCTATCAGATCACTGTTCTAGAACCTGC
CTCAGCACAGCCCAGCCGGCCCACATGCAGGCCATGAGGCAGGGGCTGCTATCACGTCACCAGCAG
GCAAAGAAAACAGCCAGACCCTCTCCAGGACGGCCTGGGGCCAAAGCGGGCTGCAGGAACTCGGCT
GGGGCACCTGAGGTTGCCCAGTCTGAGGGAGATGCCCACCCGACCCCAGGCTCCGCCCAGGCCCCA
CATTAGCACAAGCCCAGGCATGGGAGAAACAGCTGCTGAGGAAATAAAACTCCCTAAAAAAAAAAA
AAAAAAAAAAAAAAAAA

FIG. 7B.

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mMRP: 914  MYQSRPGPVAVPVQPTRPIKTFQKKNDPKDEALAKLGINGVHL-PLSTSPNQGKSSPPAV 972
           MYQSRPGPV VPVQP+RP K F +K DPKDEALAKLGING H  P   SP+ GK  PPAV
hMRP: 1    MYQSRPGPVPVPVQPSRPPKAFLRKIDPKDEALAKLGINGAHSSPPMLSPSPGKGPPPAV 60

mMRP: 973  VPRPKARPRLEPSLSIQEKQGPLRDLFGPCSPNPPTAPAPPPPPALPPPLSGEPKTPSVE 1032
            PRPKA  +L PS SI+EKQGPL DLFG       P A  PPPPPA P PL  +P T S E
hMRP: 61   APRPKAPLQLGPSSSIKEKQGPLLDLFGQ---KLPIAHTPPPPPAPPLPLPEDPGTLSAE 117

mMRP: 1033 SHALTEPMEDKNISTKLLVPSGSVCFSYANAPWKLFLRKEVFYPRENFSHPYCLSLLCQQ 1092
              LT+P+ED+ +ST+LL PSGSVCFSY   PWKLFLRKEVFYPRENFSHPY L LLC+Q
hMRP: 118  RRCLTQPVEDQGVSTQLLAPSGSVCFSYTGTPWKLFLRKEVFYPRENFSHPYYLRLLCEQ 177

mMRP: 1093 ILRDTFTESCTRISQDERHKMKGLLGDLEVSLETLDIVEDSIKKRIVVAARDNWANYFSR 1152
           ILRDTF+ESC RISQ+ER KMK LLG LEV L++L    EDS+KKRIVVAARDNWANYFSR
hMRP: 178  ILRDTFSESCIRISQNERRKMKDLLGGLEVDLDSLTTTEDSVKKRIVVAARDNWANYFSR 237

mMRP: 1153 IFPVSGESGSDVQLLGVSHRGLRLLKVTQSPSFHLDQLKTLCSYSYAEVLTVQCRGRSTL 1212
            FPVSGESGSDVQLL VSHRGLRLLKVTQ P     DQLK LCSYS+AEVL V+CRG STL
hMRP: 238  FFPVSGESGSDVQLLAVSHRGLRLLKVTQGPGLRPDQLKILCSYSFAEVLGVECRGGSTL 297

mMRP: 1213 ELSLKNEQLILHTAWARAIKAMVDLFLSELRKDSGYVIALRSYITDDNSLLSFHRGDLIR 1272
           ELSLK+EQL+LHTA ARAI+A+V+LFL+EL+KDSGYVIALRSYITD+ SLLSFHRGDLI+
hMRP: 298  ELSLKSEQLVLHTARARAIEALVELFLNELKKDSGYVIALRSYITDNCSLLSFHRGDLIK 357

mMRP: 1273 LLPVTALEPGWQFGSAGGRSGLFPDDVVQPAAAPDLSFSLGKRNSWQR 1320
           LLPV  LEPGWQFGSAGGRSGLFP D+VQPAAAPD SFS  +R+ W +
hMRP: 358  LLPVATLEPGWQFGSAGGRSGLFPADIVQPAAAPDFSFSKEQRSGWHK 405
```

Identities = 302/408 (74%), Positives = 334/408 (81%), Gaps = 4/408 (0%)

FIG. 8.

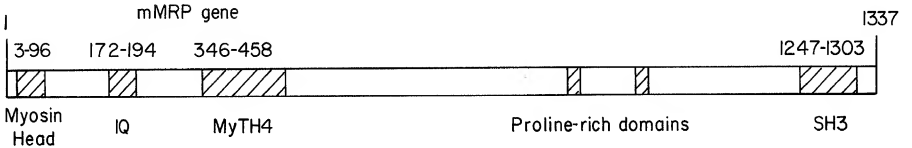

FIG. 9.

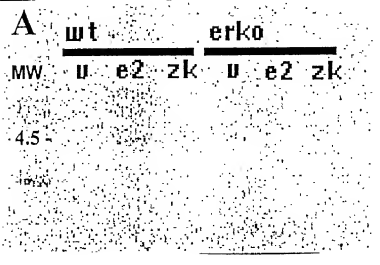

FIG. 10.

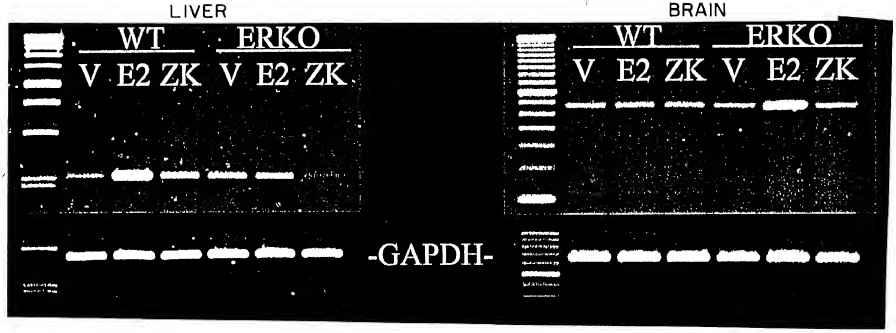

FIG. 11.

FIG. 12.

FIG. 13.